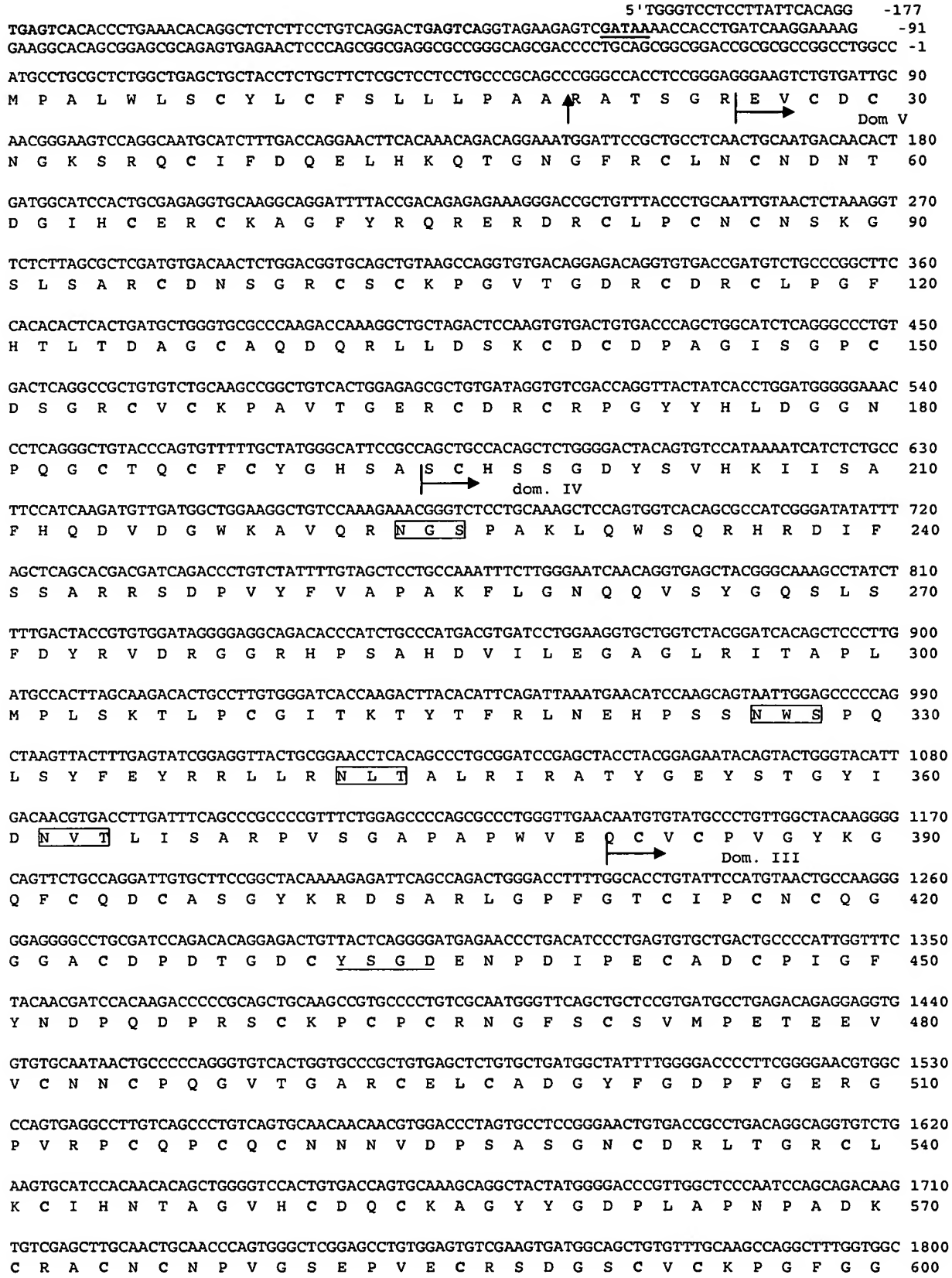
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Dom V
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D G I H C E R C K A G F Y R Q R E R D R C L P C N C N S K G 90
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S L S A R C D N S G R C S C K P G V T G D R C D R C L P G F 120
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H T L T D A G C A Q D Q R L L D S K C D C D P A G I S G P C 150
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D S G R C V C K P A V T G E R C D R C R P G Y Y H L D G G N 180
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P Q G C T Q C F C Y G H S A S C H S S G D Y S V H K I I S A 210
dom. IV
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L S Y F E Y R R L L R N L T A L R I R A T Y G E Y S T G Y I 360
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D N V T L I S A R P V S G A P A P W V E Q C V C P V G Y K G 390
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K C I H N T A G V H C D Q C K A G Y Y G D P L A P N P A D K 570
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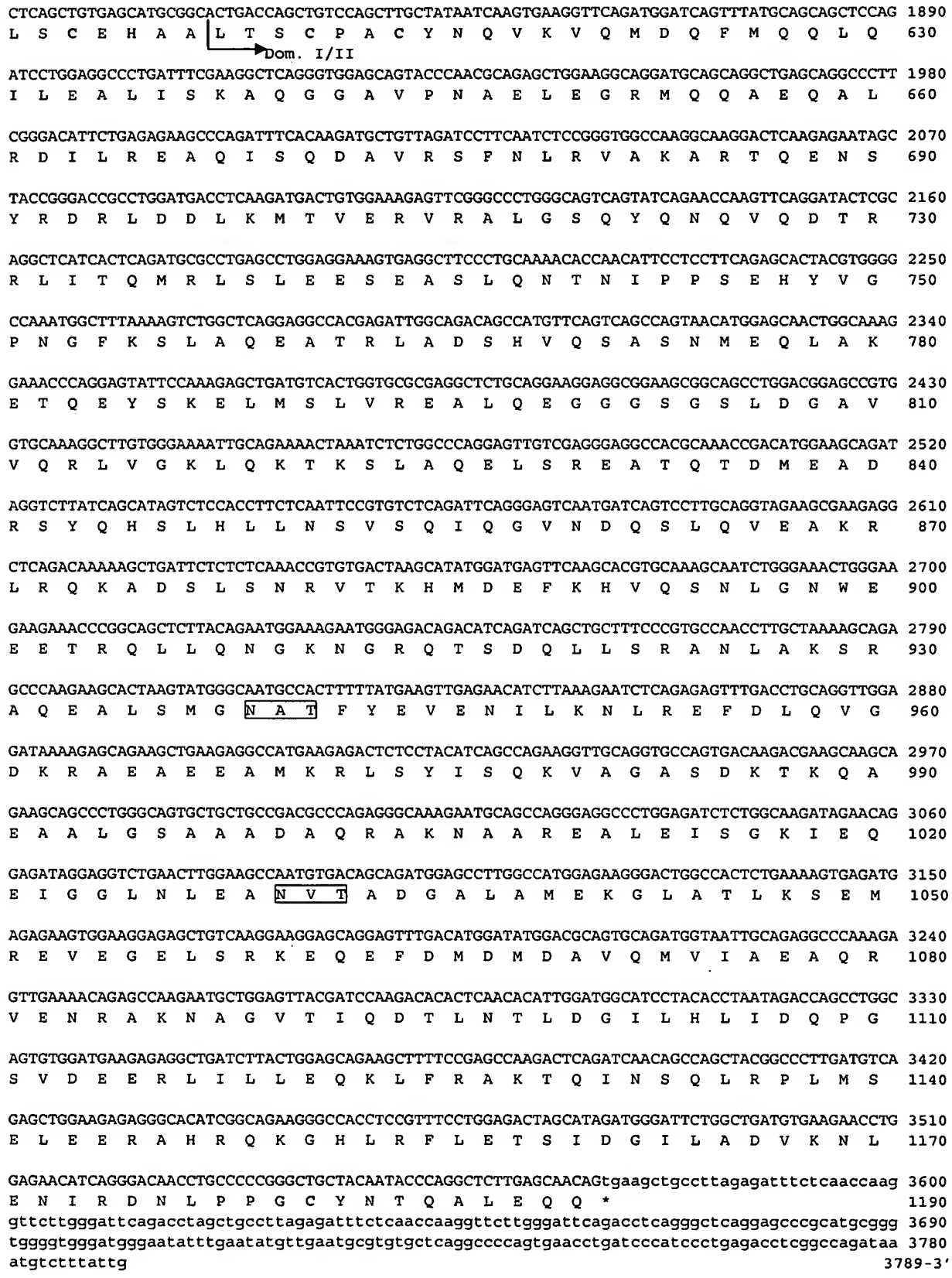
FIG. 3A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R D I L R E A Q I S Q D A V R S F N L R V A K A R T Q E N S 690
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R L I T Q M R L S L E E S E A S L Q N T N I P P S E H Y V G 750
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R E V E G E L S R K E Q E F D M D M D A V Q M V I A E A Q R 1080
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atgtctttattg 3789-3'

FIG. 3B

```
horse    1     MPALWLSCYLCFSLLLPAARATSGREVCDCNGKSRQCIFDQELHKQTGNGFRCLNCNDNTDGIHCERCKAGFYRQRERDRCLPCNCNSKGSLSARCDNSG
man      1     MPALWLGCCLCFSLLLPAARATSRREVCDCNGKSRQCIFDRELHRQTGNGFRCLNCNDNTDGIHCEKCKNGFYRHRERDRCLPCNCNSKGSLSARCDNSG
mouse    1     MPALWLSCCLGVALLLPASQATSRREVCDCNGKSRQCVFDQELHRQAGSGFRCLNCNDNTAGVHCERSREGFYQHQSKSRCLPCNCHSKGSLSAGCDNSG

horse    101   RCSCKPGVTGDRCDRCLPGFHTLTDAGCAQDQRLLDSKCDCDPAGISGPCDSGRCVCKPAVTGERCDRCRPGYYHLDGGNPQGCTQCFCYGHSASCHSSG
man      101   RCSCKPGVTGARCDRCLPGFHMLTDAGCTQDQRLLDSKCDCDPAGIAGPCDAGRCVCKPAVTGERCDRCRSGYYNLDGGNPEGCTQCFCYGHSASCRSSA
mouse    101   QCRCKPGVTGQRCDQCQPGFHMLTDAGCTRDQGQLDSKCDCDPAGISGPCDSGRCVCKPAVTGERCDRCRPRDYHLDRANPEGCTQCFCYGHSASCHASA

horse    201   DYSVHKIISAFHQDVDGWKAVQRNGSPAKLQWSQRHRDIFSSARRSDPVYFVAPAKFLGNQQVSYGQSLSFDYRVDRGGRHPSAHDVILEGAGLRITAPL
man      201   EYSVHKITSTFHQDVDGWKAVQRNGSPAKLQWSQRHQDVFSSAQRLDPVYFVAPAKFLGNQQVSYGQSLSFDYRVDRGGRHPSAHDVILEGAGLRITAPL
mouse    201   DFSVHKITSTFSQDVDGWKAVQRNGAPAKLHWSQRHRDVFSSARRSDPVYFVAPAKFLGNQQVSYGQSLSFDYRVDRGGRQPSAYDVILEGAGLQIRAPL

horse    301   MPLSKTLPCGITKTYTFRLNEHPSSNWSPQLSYFEYRRLLRNLTALR-IRATYGEYSTGYIDNVTLISARPVSGAPAPWVEQCVCPVGYKGQFCQDCASG
man      301   MPLGKTLPCGLTKTYTFRLNEHPSNNWSPQLSYFEYRRLLRNLTALR-IRATYGEYSTGYIDNVTLISARPVSGAPAPWVEQCICPVGYKGQFCQDCASG
mouse    301   MAPGKTLPCGITKTYTFRLNEHPSSHWSPQLSYFEYRRLLRNLTALLMIRATYGEYSTGYIDNVTLVSARPVLGAPAPWVERCVCLLGYKGQFCQECASG

horse    400   YKRDSARLGPFGTCIPCNCQGGGACDPDTGDCYSGDENPDIPECADCPIGFYNDPQDPRSCKPCPCRNGFSCSVMPETEEVVCNNCPQGVTGARCELCAD
man      400   YKRDSARLGPFGTCIPCNCQGGGACDPDTGDCYSGDENPDI-ECADCPIGFYNDPHDPRSCKPCPCHNGFSCSVMPETEEVVCNNCPPGVTGARCELCAD
mouse    401   YKRDSARLGAFGACVPCNCQGEGACDPDTGDCYSGDENPDI-ECADCPIGFYNDPHDPRSCKPCPCHNGFSCSVMPETEEVVCNNCPPGVTGARCELCAD

horse    500   GYFGDPFGERGPVRPCQPCQCNNNVDPSASGNCDRLTGRCLKCIHNTAGVHCDQCKAGYYGDPLAPNPADKCRACNCNPVGSEPVECRSDGSCVCKPGFG
man      499   GYFGDPFGEHGPVRPCQPCQCNNNVDPSASGNCDRLTGRCLKCIHNTAGIYCDQCKAGYFGDPLAPNPADKCRACNCNPMGSEPVGCRSDGTCVCKPGFG
mouse    500   GFFGDPFGEHGPVRPCQRCQCNNNVDPNASGNCDQLTGRCLKCIYNTAGVYCDQCKAGYFGDPLAPNPADKCRACNCSPMGAEPGECRGDGSCVCKPGFG

horse    600   GLSCEHAALTSCPACYNQVKVQMDQFMQQLQILEALISKAQGG---AVPNAELEGRMQQAEQALRDILREAQISQDAVRSFNLRVAKARTQENSYRDRLD
man      599   GPNCEHGAFS-CPACYNQVKIQMDQFMQQLQRMEALISKAQGGDG-VVPDTELEGRMQQAEQALQDILRDAQISEGASRSLGLQLAKVRSQENSYQSRLD
mouse    600   AFNCDHAALTSCPACYNQVKIQMDQFTQQLQSLEALVSKAQGGGGGGTVPVQLEGRIEQAEQALQDILGEAQISEGAMRAVAVRLAKARSQENDYKTRLD

horse    697   DLKMTVERVRALGSQYQNQVQDTRRLITQMRLSLEESEASLQNTNIPPSEHYVGPNGFKSLAQEATRLADSHVQSASNMEQLAKETQEYSKELMSLVREA
man      697   DLKMTVERVRALGSQYQNRVRDTHRLITQMQLSLAESEASLGNTNIPASDHYVGPNGFKSLAQEATRLAESHVESASNMEQLTRETEDYSKQALSLVRKA
mouse    700   DLKMTAERIRALGSQHQNRVQDTSRLISQMRLSLAGSEALLENTNIHSSEHYVGPNDFKSLAQEATRKADSHAESANAMKQLARETEDYSKQALSLARKL

horse    797   LQE—GGGSGSLDGAVVQRLVGKLQKTKSLAQELSREATQTDMEADRSYQHSLHLLNSVSQIQGVNDQSLQVEAKR-LRQKADSLSNRVTKHMDEFKHVQ
man      797   LHEGVGSGSGSPDGAVVQGLVEKLEKTKSLAQQLTREATQAEIEADRSYQHSLRLLDSVSRLQGVSDQSFQVEEAKRIKQKADSLSTLVTRHMDEFKRTQ
mouse    800   LSG—GGGSGSWDSSVVQGLMGKLEKTKSLSQQLSLEGTQADIEADRSYQHSLRLLDSASQLQGVSDLSFQVEAKR-IRQKADSLSNLVTRQTDAFTRVR

horse    894   SNLGNWEEETRQLLQNGKNGRQTSDQLLSRANLAKSRAQEALSMGNATFYEVENILKNLREFDLQVGDKRAEAEEAMKRLSYISQKVAGASDKTKQAEAA
man      897   KNLGNWKEEAQQLLQNGKSGREKSDQLLSRANLAKSRAQEALSMGNATFYEVESILKNLREFDLQVDNRKAEAEEAMKRLSYISQKVSDASDKTQQAERA
mouse    897   NNLGNWEKETRQLLQTGKDRRQTSDQLLSRANLAKNRAQEALSMGNATFYEVENILKNLREFDLQVEDRKAEAEEAMKRLSSISQKVADASDKTQQAETA

horse    994   LGSAAADAQRAKNAAREALEISGKIEQEIGGLNLEANVTADGALAMEKGLATLKSEMREVEGELSRKEQEFDMDMDAVQMVIAEAQRVENRAKNAGVTIQ
man      997   LGSAAADAQRAKNGAGEALEISSEIEQEIGSLNLEANVTADGALAMEKGLASLKSEMREVEGELERKELEFDTNMDAVQMVITEAQKVDTRAKNAGVTIQ
mouse    997   LGSATADTQRAKNAAREALEISSEIELEIGSLNLEANVTADGALAMEKGTATLKSEMREMI-ELARKELEFDTDKDTVQLVITEAQQADARATSAGVTIQ

horse    1094  DTLNTLDGILHLIDQPGSVDEERLILLEQKLFRAKTQINSQLRPLMSELEERAHRQKGHLRFLETSIDGILADVKNLENIRDNLPPGCYNTQALEQQ
man      1097  DTLNTLDGLLHLMDQPLSVDEEGLVLLEQKLSRAKTQINSQLRPMMSELEERARQQRGHLHLLETSIDGILADVKNLENIRDNLPPGCYNTQALEQQ
mouse    1096  DTLNTLDGILHLIDQPGSVDEEGMMLLEQGLFQAKTQINSRLRPLMSDLEERVRRQRNHLHLLETSIDGILADVKNLENIRDNLPPGCYNTQALEQQ
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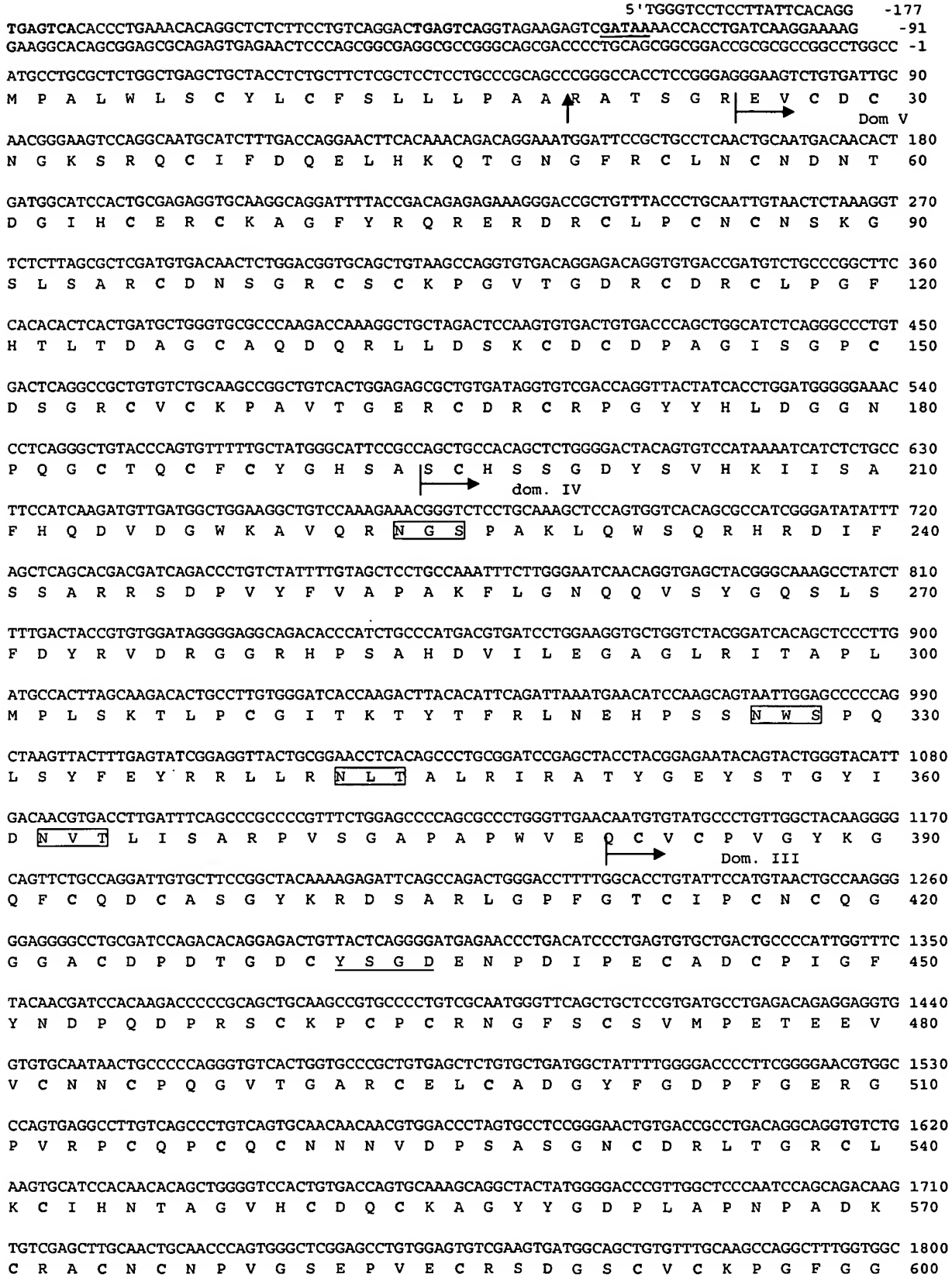
FIG. 4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M P A L W L S C Y L C F S L L L P A A R A T S G R E V C D C 30
Dom V
AACGGGAAGTCCAGGCAATGCATCTTTGACCAGGAACTTCACAAACAGACAGGAAATGGATTCCGCTGCCTCAACTGCAATGACAACACT 180
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D G I H C E R C K A G F Y R Q R E R D R C L P C N C N S K G 90
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S L S A R C D N S G R C S C K P G V T G D R C D R C L P G F 120
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D S G R C V C K P A V T G E R C D R C R P G Y Y H L D G G N 180
CCTCAGGGCTGTACCCAGTGTTTTTGCTATGGGCATTCCGCCAGCTGCCACAGCTCTGGGGACTACAGTGTCCATAAAATCATCTCTGCC 630
P Q G C T Q C F C Y G H S A S C H S S G D Y S V H K I I S A 210
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S S A R R S D P V Y F V A P A K F L G N Q Q V S Y G Q S L S 270
TTTGACTACCGTGTGGATAGGGGAGGCAGACACCCATCTGCCCATGACGTGATCCTGGAAGGTGCTGGTCTACGGATCACAGCTCCCTTG 900
F D Y R V D R G G R H P S A H D V I L E G A G L R I T A P L 300
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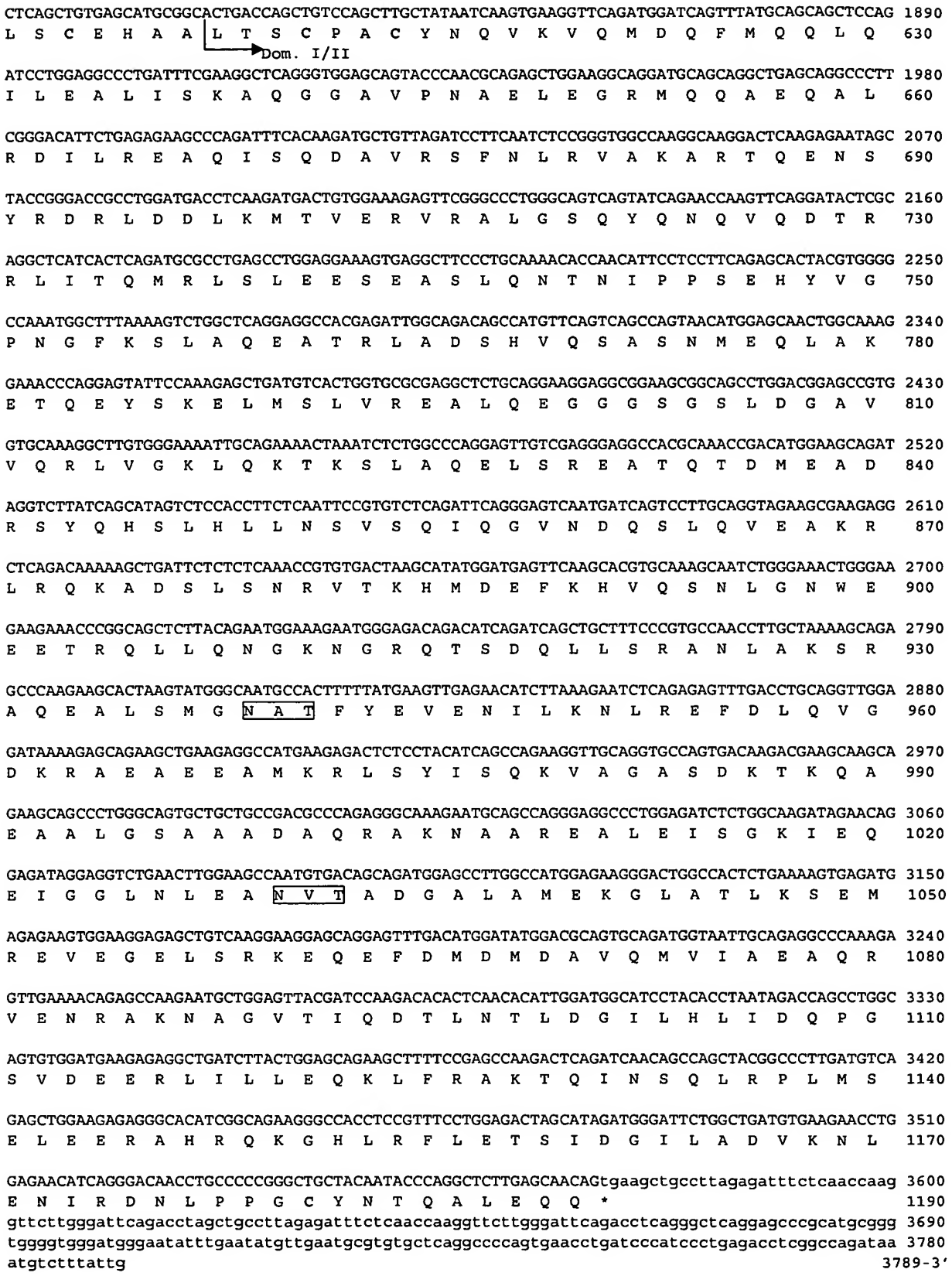
FIG. 3A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TACCGGGACCGCCTGGATGACCTCAAGATGACTGTGGAAAGAGTTCGGGCCCTGGGCAGTCAGTATCAGAACCAAGTTCAGGATACTCGC 2160
Y R D R L D D L K M T V E R V R A L G S Q Y Q N Q V Q D T R 730
AGGCTCATCACTCAGATGCGCCTGAGCCTGGAGGAAAGTGAGGCTTCCCTGCAAAACACCAACATTCCTCCTTCAGAGCACTACGTGGGG 2250
R L I T Q M R L S L E E S E A S L Q N T N I P P S E H Y V G 750
CCAAATGGCTTTAAAAGTCTGGCTCAGGAGGCCACGAGATTGGCAGACAGCCATGTTCAGTCAGCCAGTAACATGGAGCAACTGGCAAAG 2340
P N G F K S L A Q E A T R L A D S H V Q S A S N M E Q L A K 780
GAAACCCAGGAGTATTCCAAAGAGCTGATGTCACTGGTGCGCGAGGCTCTGCAGGAAGGAGGCGGAAGCGGCAGCCTGGACGGAGCCGTG 2430
E T Q E Y S K E L M S L V R E A L Q E G G G S G S L D G A V 810
GTGCAAAGGCTTGTGGGAAAATTGCAGAAAACTAAATCTCTGGCCCAGGAGTTGTCGAGGGAGGCCACGCAAACCGACATGGAAGCAGAT 2520
V Q R L V G K L Q K T K S L A Q E L S R E A T Q T D M E A D 840
AGGTCTTATCAGCATAGTCTCCACCTTCTCAATTCCGTGTCTCAGATTCAGGGAGTCAATGATCAGTCCTTGCAGGTAGAAGCGAAGAGG 2610
R S Y Q H S L H L L N S V S Q I Q G V N D Q S L Q V E A K R 870
CTCAGACAAAAAGCTGATTCTCTCTCAAACCGTGTGACTAAGCATATGGATGAGTTCAAGCACGTGCAAAGCAATCTGGGAAACTGGGAA 2700
L R Q K A D S L S N R V T K H M D E F K H V Q S N L G N W E 900
GAAGAAACCCGGCAGCTCTTACAGAATGGAAAGAATGGGAGACAGACATCAGATCAGCTGCTTTCCCGTGCCAACCTTGCTAAAAGCAGA 2790
E E T R Q L L Q N G K N G R Q T S D Q L L S R A N L A K S R 930
GCCCAAGAAGCACTAAGTATGGGCAATGCCACTTTTTATGAAGTTGAGAACATCTTAAAGAATCTCAGAGAGTTTGACCTGCAGGTTGGA 2880
A Q E A L S M G N A T F Y E V E N I L K N L R E F D L Q V G 960
GATAAAAGAGCAGAAGCTGAAGAGGCCATGAAGAGACTCTCCTACATCAGCCAGAAGGTTGCAGGTGCCAGTGACAAGACGAAGCAAGCA 2970
D K R A E A E E A M K R L S Y I S Q K V A G A S D K T K Q A 990
GAAGCAGCCCTGGGCAGTGCTGCTGCCGACGCCCAGAGGGCAAAGAATGCAGCCAGGGAGGCCCTGGAGATCTCTGGCAAGATAGAACAG 3060
E A A L G S A A A D A Q R A K N A A R E A L E I S G K I E Q 1020
GAGATAGGAGGTCTGAACTTGGAAGCCAATGTGACAGCAGATGGAGCCTTGGCCATGGAGAAGGGACTGGCCACTCTGAAAAGTGAGATG 3150
E I G G L N L E A N V T A D G A L A M E K G L A T L K S E M 1050
AGAGAAGTGGAAGGAGAGCTGTCAAGGAAGGAGCAGGAGTTTGACATGGATATGGACGCAGTGCAGATGGTAATTGCAGAGGCCCAAAGA 3240
R E V E G E L S R K E Q E F D M D M D A V Q M V I A E A Q R 1080
GTTGAAAACAGAGCCAAGAATGCTGGAGTTACGATCCAAGACACACTCAACACATTGGATGGCATCCTACACCTAATAGACCAGCCTGGC 3330
V E N R A K N A G V T I Q D T L N T L D G I L H L I D Q P G 1110
AGTGTGGATGAAGAGAGGCTGATCTTACTGGAGCAGAAGCTTTTCCGAGCCAAGACTCAGATCAACAGCCAGCTACGGCCCTTGATGTCA 3420
S V D E E R L I L L E Q K L F R A K T Q I N S Q L R P L M S 1140
GAGCTGGAAGAGAGGGCACATCGGCAGAAGGGCCACCTCCGTTTCCTGGAGACTAGCATAGATGGGATTCTGGCTGATGTGAAGAACCTG 3510
E L E E R A H R Q K G H L R F L E T S I D G I L A D V K N L 1170
GAGAACATCAGGGACAACCTGCCCCCGGGCTGCTACAATACCCAGGCTCTTGAGCAACAGtgaagctgccttagagatttctcaaccaag 3600
E N I R D N L P P G C Y N T Q A L E Q Q * 1190
gttcttgggattcagacctagctgccttagagatttctcaaccaaggttcttgggattcagacctcagggctcaggagcccgcatgcggg 3690
tggggtgggatgggaatatttgaatatgttgaatgcgtgtgctcaggccccagtgaacctgatcccatccctgagacctcggccagataa 3780
atgtctttattg 3789-3'

FIG. 3B

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horse    1     MPALWLSCYLCFSLLLPAARATSGREVCDCNGKSRQCIFDQELHKQTGNGFRCLNCNDNTDGIHCERCKAGFYRQRERDRCLPCNCNSKGSLSARCDNSG
man      1     MPALWLGCCLCFSLLLPAARATSRREVCDCNGKSRQCIFDRELHRQTGNGFRCLNCNDNTDGIHCEKCKNGFYRHRERDRCLPCNCNSKGSLSARCDNSG
mouse    1     MPALWLSCCLGVALLLPASQATSRREVCDCNGKSRQCVFDQELHRQAGSGFRCLNCNDNTAGVHCERSREGFYQHQSKSRCLPCNCHSKGSLSAGCDNSG

horse    101   RCSCKPGVTGDRCDRCLPGFHTLTDAGCAQDQRLLDSKCDCDPAGISGPCDSGRCVCKPAVTGERCDRCRPGYYHLDGGNPQGCTQCFCYGHSASCHSSG
man      101   RCSCKPGVTGARCDRCLPGFHMLTDAGCTQDQRLLDSKCDCDPAGIAGPCDAGRCVCKPAVTGERCDRCRSGYYNLDGGNPEGCTQCFCYGHSASCRSSA
mouse    101   QCRCKPGVTGQRCDQCQPGFHMLTDAGCTRDQGQLDSKCDCDPAGISGPCDSGRCVCKPAVTGERCDRCRPRDYHLDRANPEGCTQCFCYGHSASCHASA

horse    201   DYSVHKIISAFHQDVDGWKAVQRNGSPAKLQWSQRHRDIFSSARRSDPVYFVAPAKFLGNQQVSYGQSLSFDYRVDRGGRHPSAHDVILEGAGLRITAPL
man      201   EYSVHKITSTFHQDVDGWKAVQRNGSPAKLQWSQRHQDVFSSAQRLDPVYFVAPAKFLGNQQVSYGQSLSFDYRVDRGGRHPSAHDVILEGAGLRITAPL
mouse    201   DFSVHKITSTFSQDVDGWKAVQRNGAPAKLHWSQRHRDVFSSARRSDPVYFVAPAKFLGNQQVSYGQSLSFDYRVDRGGRQPSAYDVILEGAGLQIRAPL

horse    301   MPLSKTLPCGITKTYTFRLNEHPSSNWSPQLSYFEYRRLLRNLTALR-IRATYGEYSTGYIDNVTLISARPVSGAPAPWVEQCVCPVGYKGQFCQDCASG
man      301   MPLGKTLPCGLTKTYTFRLNEHPSNNWSPQLSYFEYRRLLRNLTALR-IRATYGEYSTGYIDNVTLISARPVSGAPAPWVEQCICPVGYKGQFCQDCASG
mouse    301   MAPGKTLPCGITKTYTFRLNEHPSSHWSPQLSYFEYRRLLRNLTALLMIRATYGEYSTGYIDNVTLVSARPVLGAPAPWVERCVCLLGYKGQFCQECASG

horse    400   YKRDSARLGPFGTCIPCNCQGGGACDPDTGDCYSGDENPDIPECADCPIGFYNDPQDPRSCKPCPCRNGFSCSVMPETEEVVCNNCPQGVTGARCELCAD
man      400   YKRDSARLGPFGTCIPCNCQGGGACDPDTGDCYSGDENPDI-ECADCPIGFYNDPHDPRSCKPCPCHNGFSCSVMPETEEVVCNNCPPGVTGARCELCAD
mouse    401   YKRDSARLGAFGACVPCNCQGEGACDPDTGDCYSGDENPDI-ECADCPIGFYNDPHDPRSCKPCPCHNGFSCSVMPETEEVVCNNCPPGVTGARCELCAD

horse    500   GYFGDPFGERGPVRPCQPCQCNNNVDPSASGNCDRLTGRCLKCIHNTAGVHCDQCKAGYYGDPLAPNPADKCRACNCNPVGSEPVECRSDGSCVCKPGFG
man      499   GYFGDPFGEHGPVRPCQPCQCNNNVDPSASGNCDRLTGRCLKCIHNTAGIYCDQCKAGYFGDPLAPNPADKCRACNCNPMGSEPVGCRSDGTCVCKPGFG
mouse    500   GFFGDPFGEHGPVRPCQRCQCNNNVDPNASGNCDQLTGRCLKCIYNTAGVYCDQCKAGYFGDPLAPNPADKCRACNCSPMGAEPGECRGDGSCVCKPGFG

horse    600   GLSCEHAALTSCPACYNQVKVQMDQFMQQLQILEALISKAQGG---AVPNAELEGRMQQAEQALRDILREAQISQDAVRSFNLRVAKARTQENSYRDRLD
man      599   GPNCEHGAFS-CPACYNQVKIQMDQFMQQLQRMEALISKAQGGDG-VVPDTELEGRMQQAEQALQDILRDAQISEGASRSLGLQLAKVRSQENSYQSRLD
mouse    600   AFNCDHAALTSCPACYNQVKIQMDQFTQQLQSLEALVSKAQGGGGGGTVPVQLEGRIEQAEQALQDILGEAQISEGAMRAVAVRLAKARSQENDYKTRLD

horse    697   DLKMTVERVRALGSQYQNQVQDTRRLITQMRLSLEESEASLQNTNIPPSEHYVGPNGFKSLAQEATRLADSHVQSASNMEQLAKETQEYSKELMSLVREA
man      697   DLKMTVERVRALGSQYQNRVRDTHRLITQMQLSLAESEASLGNTNIPASDHYVGPNGFKSLAQEATRLAESHVESASNMEQLTRETEDYSKQALSLVRKA
mouse    700   DLKMTAERIRALGSQHQNRVQDTSRLISQMRLSLAGSEALLENTNIHSSEHYVGPNDFKSLAQEATRKADSHAESANAMKQLARETEDYSKQALSLARKL

horse    797   LQE—GGGSGSLDGAVVQRLVGKLQKTKSLAQELSREATQTDMEADRSYQHSLHLLNSVSQIQGVNDQSLQVEAKR-LRQKADSLSNRVTKHMDEFKHVQ
man      797   LHEGVGSGSGSPDGAVVQGLVEKLEKTKSLAQQLTREATQAEIEADRSYQHSLRLLDSVSRLQGVSDQSFQVEEAKRIKQKADSLSTLVTRHMDEFKRTQ
mouse    800   LSG—GGGSGSWDSSVVQGLMGKLEKTKSLSQQLSLEGTQADIEADRSYQHSLRLLDSASQLQGVSDLSFQVEAKR-IRQKADSLSNLVTRQTDAFTRVR

horse    894   SNLGNWEEETRQLLQNGKNGRQTSDQLLSRANLAKSRAQEALSMGNATFYEVENILKNLREFDLQVGDKRAEAEEAMKRLSYISQKVAGASDKTKQAEAA
man      897   KNLGNWKEEAQQLLQNGKSGREKSDQLLSRANLAKSRAQEALSMGNATFYEVESILKNLREFDLQVDNRKAEAEEAMKRLSYISQKVSDASDKTQQAERA
mouse    897   NNLGNWEKETRQLLQTGKDRRQTSDQLLSRANLAKNRAQEALSMGNATFYEVENILKNLREFDLQVEDRKAEAEEAMKRLSSISQKVADASDKTQQAETA

horse    994   LGSAAADAQRAKNAAREALEISGKIEQEIGGLNLEANVTADGALAMEKGLATLKSEMREVEGELSRKEQEFDMDMDAVQMVIAEAQRVENRAKNAGVTIQ
man      997   LGSAAADAQRAKNGAGEALEISSEIEQEIGSLNLEANVTADGALAMEKGLASLKSEMREVEGELERKELEFDTNMDAVQMVITEAQKVDTRAKNAGVTIQ
mouse    997   LGSATADTQRAKNAAREALEISSEIELEIGSLNLEANVTADGALAMEKGTATLKSEMREMI-ELARKELEFDTDKDTVQLVITEAQQADARATSAGVTIQ

horse    1094  DTLNTLDGILHLIDQPGSVDEERLILLEQKLFRAKTQINSQLRPLMSELEERAHRQKGHLRFLETSIDGILADVKNLENIRDNLPPGCYNTQALEQQ
man      1097  DTLNTLDGLLHLMDQPLSVDEEGLVLLEQKLSRAKTQINSQLRPMMSELEERARQQRGHLHLLETSIDGILADVKNLENIRDNLPPGCYNTQALEQQ
mouse    1096  DTLNTLDGILHLIDQPGSVDEEGMMLLEQGLFQAKTQINSRLRPLMSDLEERVRRQRNHLHLLETSIDGILADVKNLENIRDNLPPGCYNTQALEQQ
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FIG. 4